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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/833,245

DATE: 04/30/2001
TIME: 09:10:20

Input Set : N:\jumbos\PF546PCTSL.txt
Output Set: N:\CRF3\04302001\I833245.raw

4 <110> APPLICANT: Human Genome Sciences, Inc.
 6 <120> TITLE OF INVENTION: Albumin Fusion Proteins
 8 <130> FILE REFERENCE: PF546PCT
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/833,245
 11 <141> CURRENT FILING DATE: 2001-04-12
 13 <150> PRIOR APPLICATION NUMBER: 60/229, 358
 14 <151> PRIOR FILING DATE: 2000-04-12
 16 <150> PRIOR APPLICATION NUMBER: 60/256, 931
 17 <151> PRIOR FILING DATE: 2000-12-21
 19 <150> PRIOR APPLICATION NUMBER: 60/199, 384
 20 <151> PRIOR FILING DATE: 2000-04-25
 22 <160> NUMBER OF SEQ ID NOS: 2267
 24 <170> SOFTWARE: PatentIn Ver. 2.1
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 23
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Artificial Sequence
 31 <220> FEATURE:
 32 <221> NAME/KEY: primer_bind
 33 <223> OTHER INFORMATION: primer useful to clone human growth hormone cDNA
 35 <400> SEQUENCE: 1
 36 cccaaaggatt cccttatcca ggc 23
 39 <210> SEQ ID NO: 2
 40 <211> LENGTH: 33
 41 <212> TYPE: DNA
 42 <213> ORGANISM: Artificial Sequence
 44 <220> FEATURE:
 45 <221> NAME/KEY: primer_bind
 46 <223> OTHER INFORMATION: primer useful to clone human growth hormone cDNA
 48 <400> SEQUENCE: 2
 49 gggaagctta gaagccacag gatccctcca cag 33
 52 <210> SEQ ID NO: 3
 53 <211> LENGTH: 16
 54 <212> TYPE: DNA
 55 <213> ORGANISM: Artificial Sequence
 57 <220> FEATURE:
 58 <221> NAME/KEY: misc_structure
 59 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA fragments
 60 with non-cohesive ends.
 62 <400> SEQUENCE: 3
 63 gataaaaggatt cccaaac 16
 66 <210> SEQ ID NO: 4
 67 <211> LENGTH: 17
 68 <212> TYPE: DNA
 69 <213> ORGANISM: Artificial Sequence
 71 <220> FEATURE:
 72 <221> NAME/KEY: misc_structure

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73 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA fragments
74 with non-cohesive ends.
76 <400> SEQUENCE: 4
77 aattgttggg aatcttt
80 <210> SEQ ID NO: 5
81 <211> LENGTH: 17
82 <212> TYPE: DNA
83 <213> ORGANISM: Artificial Sequence
85 <220> FEATURE:
86 <221> NAME/KEY: misc_structure
87 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA fragments
88 with non-cohesive ends.
90 <400> SEQUENCE: 5
91 ttaggcttat tcccaac
94 <210> SEQ ID NO: 6
95 <211> LENGTH: 18
96 <212> TYPE: DNA
97 <213> ORGANISM: Artificial Sequence
99 <220> FEATURE:
100 <221> NAME/KEY: misc_structure
101 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA fragments
102 with non-cohesive ends.
104 <400> SEQUENCE: 6
105 aattgttggg aataagcc
108 <210> SEQ ID NO: 7
109 <211> LENGTH: 24
110 <212> TYPE: PRT
111 <213> ORGANISM: Artificial Sequence
113 <220> FEATURE:
114 <221> NAME/KEY: SITE
115 <222> LOCATION: 1)...(19)
116 <223> OTHER INFORMATION: invertase leader sequence
118 <220> FEATURE:
119 <221> NAME/KEY: SITE
120 <222> LOCATION: 20)...(24)
121 <223> OTHER INFORMATION: first 5 amino acids of mature human serum albumin
123 <400> SEQUENCE: 7
124 Met Leu Leu Gln Ala Phe Leu Phe Leu Leu Ala Gly Phe Ala Ala Lys
125 1 5 10 15
127 Ile Ser Ala Asp Ala His Lys Ser
128 20
131 <210> SEQ ID NO: 8
132 <211> LENGTH: 21
133 <212> TYPE: DNA
134 <213> ORGANISM: Artificial Sequence
136 <220> FEATURE:
137 <221> NAME/KEY: misc_structure
138 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA
139 fragments with non-cohesive ends.

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141 <400> SEQUENCE: 8 21
142 gagatgcaca cctgagttag g
145 <210> SEQ ID NO: 9
146 <211> LENGTH: 27
147 <212> TYPE: DNA
148 <213> ORGANISM: Artificial Sequence
150 <220> FEATURE:
151 <221> NAME/KEY: misc_structure
152 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA
fragments with non-cohesive ends.
155 <400> SEQUENCE: 9 27
156 gatcctgtgg cttagatgca cacaaga
159 <210> SEQ ID NO: 10
160 <211> LENGTH: 24
161 <212> TYPE: DNA
162 <213> ORGANISM: Artificial Sequence
164 <220> FEATURE:
165 <221> NAME/KEY: misc_structure
166 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA
fragments with non-cohesive ends.
169 <400> SEQUENCE: 10 24
170 ctcttggatg catcgaaagcc acag
173 <210> SEQ ID NO: 11
174 <211> LENGTH: 30
175 <212> TYPE: DNA
176 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <221> NAME/KEY: misc_structure
180 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA
fragments with non-cohesive ends.
183 <400> SEQUENCE: 11 30
184 tgtgaaagag cctcagaatt tattcccaac
187 <210> SEQ ID NO: 12
188 <211> LENGTH: 31
189 <212> TYPE: DNA
190 <213> ORGANISM: Artificial Sequence
192 <220> FEATURE:
193 <221> NAME/KEY: misc_structure
194 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA
fragments with non-cohesive ends.
197 <400> SEQUENCE: 12 31
198 aattgttggg aataaaattct gaggcttttc c
201 <210> SEQ ID NO: 13
202 <211> LENGTH: 47
203 <212> TYPE: DNA
204 <213> ORGANISM: Artificial Sequence
206 <220> FEATURE:
207 <221> NAME/KEY: misc_structure
208 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA

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209 fragments with non-cohesive ends.
211 <400> SEQUENCE: 13
212 ttaggcttag gtggcggtgg atccggcggt ggtggatctt tcccaac 47
215 <210> SEQ ID NO: 14
216 <211> LENGTH: 48
217 <212> TYPE: DNA
218 <213> ORGANISM: Artificial Sequence
220 <220> FEATURE:
221 <221> NAME/KEY: misc_structure
222 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA
223 fragments with non-cohesive ends.
225 <400> SEQUENCE: 14
226 aattgttggg aaagatccac caccgcggga tccaccgcga cctaaggcc 48
229 <210> SEQ ID NO: 15
230 <211> LENGTH: 62
231 <212> TYPE: DNA
232 <213> ORGANISM: Artificial Sequence
234 <220> FEATURE:
235 <221> NAME/KEY: misc_structure
236 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA
237 fragments with non-cohesive ends.
239 <400> SEQUENCE: 15
240 ttaggcttag gcgggtggatcttggc ggcggatctg gtggcggtgg atcctccca 60 62
241 ac
244 <210> SEQ ID NO: 16
245 <211> LENGTH: 63
246 <212> TYPE: DNA
247 <213> ORGANISM: Artificial Sequence
249 <220> FEATURE:
250 <221> NAME/KEY: misc_structure
251 <223> OTHER INFORMATION: synthetic oligonucleotide used to join DNA
252 fragments with non-cohesive ends.
254 <400> SEQUENCE: 16
255 aattgttggg aaggatccac cgccaccaga tccggcgcga ccagatccac caccgcctaa 60 63
256 gcc
259 <210> SEQ ID NO: 17
260 <211> LENGTH: 1782
261 <212> TYPE: DNA
262 <213> ORGANISM: Homo sapiens
264 <220> FEATURE:
265 <221> NAME/KEY: CDS
266 <222> LOCATION: (1)..(1755)
269 <400> SEQUENCE: 17
270 gat gca cac aag agt gag gtt gct cat cgg ttt aaa gat ttg gga gaa 48
271 Asp Ala His Lys Ser Glu Val Ala His Arg Phe Lys Asp Leu Gly Glu
272 1 5 10 15
274 gaa aat ttc aaa gcc ttg gtg ttg att gcc ttt gct cag tat ctt cag 96
275 Glu Asn Phe Lys Ala Leu Val Leu Ile Ala Phe Ala Gln Tyr Leu Gln
276 20 25 30

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278 cag tgt cca ttt gaa gat cat gta aaa tta gtg aat gaa gta act gaa	144
279 Gln Cys Pro Phe Glu Asp His Val Lys Leu Val Asn Glu Val Thr Glu	
280 35 40 45	
282 ttt gca aaa aca tgt gtt gct gat gag tca gct gaa aat tgt gac aaa	192
283 Phe Ala Lys Thr Cys Val Ala Asp Glu Ser Ala Glu Asn Cys Asp Lys	
284 50 55 60	
286 tca ctt cat acc ctt ttt gga gac aaa tta tgc aca gtt gca act ctt	240
287 Ser Leu His Thr Leu Phe Gly Asp Lys Leu Cys Thr Val Ala Thr Leu	
288 65 70 75 80	
290 cgt gaa acc tat ggt gaa atg gct gac tgc tgt gca aaa caa gaa cct	288
291 Arg Glu Thr Tyr Gly Glu Met Ala Asp Cys Cys Ala Lys Gln Glu Pro	
292 85 90 95	
294 gag aga aat gaa tgc ttc ttg caa cac aaa gat gac aac cca aac ctc	336
295 Glu Arg Asn Glu Cys Phe Leu Gln His Lys Asp Asp Asn Pro Asn Leu	
296 100 105 110	
298 ccc cga ttg gtg aga cca gag gtt gat gtg atg tgc act gct ttt cat	384
299 Pro Arg Leu Val Arg Pro Glu Val Asp Val Met Cys Thr Ala Phe His	
300 115 120 125	
302 gac aat gaa gag aca ttt ttg aaa aaa tac tta tat gaa att gcc aga	432
303 Asp Asn Glu Glu Thr Phe Leu Lys Lys Tyr Leu Tyr Glu Ile Ala Arg	
304 130 135 140	
306 aga cat cct tac ttt tat gcc ccg gaa ctc ctt ttc ttt gct aaa agg	480
307 Arg His Pro Tyr Phe Tyr Ala Pro Glu Leu Leu Phe Ala Lys Arg	
308 145 150 155 160	
310 tat aaa gct gct ttt aca gaa tgt tgc caa gct gct gat aaa gct gcc	528
311 Tyr Lys Ala Ala Phe Thr Glu Cys Cys Gln Ala Ala Asp Lys Ala Ala	
312 165 170 175	
314 tgc ctg ttg cca aag ctc gat gaa ctt ccg gat gaa ggg aag gct tcg	576
315 Cys Leu Leu Pro Lys Leu Asp Glu Leu Arg Asp Glu Gly Lys Ala Ser	
316 180 185 190	
318 tct gcc aaa cag aga ctc aaa tgt gcc agt ctc caa aaa ttt gga gaa	624
319 Ser Ala Lys Gln Arg Leu Lys Cys Ala Ser Leu Gln Lys Phe Gly Glu	
320 195 200 205	
322 aga gct ttc aaa gca tgg gca gtg gct cgc ctg agc cag aga ttt ccc	672
323 Arg Ala Phe Lys Ala Trp Ala Val Ala Arg Leu Ser Gln Arg Phe Pro	
324 210 215 220	
326 aaa gct gag ttt gca gaa gtt tcc aag tta gtg aca gat ctt acc aaa	720
327 Lys Ala Glu Phe Ala Glu Val Ser Lys Leu Val Thr Asp Leu Thr Lys	
328 225 230 235 240	
330 gtc cac acg gaa tgc tgc cat gga gat ctg ctt gaa tgt gct gat gac	768
331 Val His Thr Glu Cys Cys His Gly Asp Leu Leu Glu Cys Ala Asp Asp	
332 245 250 255	
334 agg gcg gac ctt gcc aag tat atc tgt gaa aat cag gat tcg atc tcc	816
335 Arg Ala Asp Leu Ala Lys Tyr Ile Cys Glu Asn Gln Asp Ser Ile Ser	
336 260 265 270	
338 agt aaa ctg aag gaa tgc tgt gaa aaa cct ctg ttg gaa aaa tcc cac	864
339 Ser Lys Leu Lys Glu Cys Cys Glu Lys Pro Leu Leu Glu Lys Ser His	
340 275 280 285	
342 tgc att gcc gaa gtg gaa aat gat gag atg cct gct gac ttg cct tca	912

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields for each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
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Input Set : N:\jumbos\PF546PCTSL.txt
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L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:708 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:798 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:887 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:977 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:1287 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:1854 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74
L:1908 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:76
L:1952 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77
L:1955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77
L:1987 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:1990 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:1993 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:2304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82
L:2307 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82
L:2310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82
L:2356 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83
L:2359 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83
L:2362 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83
L:2466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86
L:2497 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:87
L:2557 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:89
L:2596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:91
L:2693 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95
L:2699 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95
L:2720 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95
L:2823 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99
L:2992 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:101
L:2995 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:101
L:3198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:104
L:3201 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:104
L:3204 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:104
L:3514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:115
L:3523 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:115
L:3702 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:124
L:3705 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:124
L:3720 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:124
L:3759 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:125
L:3901 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127
L:3904 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127
L:3916 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127
L:4013 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:4177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:137
L:4534 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:146
L:4543 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:146
L:4549 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:146
L:4555 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:146

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L:4567 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:146
L:4787 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154
L:4790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:154